



## **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 101028,075A  
Source: OIP  
Date Processed by STIC: 10/30/02

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

# Raw Sequence Listing Error Summary

01PE

## ERROR DETECTED

## SUGGESTED CORRECTION

SERIAL NUMBER: 10/028,075A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1      Wrapped Nucleics  
    Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2      Invalid Line Length      The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3      Misaligned Amino  
    Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4      Non-ASCII      The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5      Variable Length      Sequence(s)          contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6      PatentIn 2.0  
    "bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)         . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7      Skipped Sequences  
    (OLD RULES)      Sequence(s)          missing. If intentional, please insert the following lines for each skipped sequence:  
    (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
    (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
    (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
    This sequence is intentionally skipped  
  
    Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8      Skipped Sequences  
    (NEW RULES)      Sequence(s)          missing. If intentional, please insert the following lines for each skipped sequence.  
    <210> sequence id number  
    <400> sequence id number  
    000
- 9      Use of n's or Xaa's  
    (NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
    Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
    In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10      Invalid <213>  
    Response      Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11      Use of <220>      Sequence(s)          missing the <220> "Feature" and associated numeric identifiers and responses.  
    Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
    (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12      PatentIn 2.0  
    "bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13      Misuse of n      n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



Does Not Comply  
Corrected Diskette Needed

OIPE

## RAW SEQUENCE LISTING

DATE: 10/30/2002

PATENT APPLICATION: US/10/028,075A

TIME: 15:01:56

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

```

3 <110> APPLICANT: Khan, Nisar A.
4   Benner, Robert
6 <120> TITLE OF INVENTION: Gene regulator
8 <130> FILE REFERENCE: 2183-5223US
10 <140> CURRENT APPLICATION NUMBER: 10/028,075A
11 <141> CURRENT FILING DATE: 2001-12-21
13 <150> PRIOR APPLICATION NUMBER: EP 01203748.7
14 <151> PRIOR FILING DATE: 2001-10-04
16 <160> NUMBER OF SEQ ID NOS: 175
18 <170> SOFTWARE: PatentIn Ver. 2.1
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 4
22 <212> TYPE: PRT
23 <213> ORGANISM: Artificial Sequence
25 <220> FEATURE:
26 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide
28 <400> SEQUENCE: 1
29 Leu Gln Gly Val
30   1
32 <210> SEQ ID NO: 2
33 <211> LENGTH: 4
34 <212> TYPE: PRT
35 <213> ORGANISM: Artificial Sequence
37 <220> FEATURE:
38 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide
40 <400> SEQUENCE: 2
41 Ala Gln Gly Val
42   1
43 <210> SEQ ID NO: 3
44 <211> LENGTH: 6
45 <212> TYPE: PRT
46 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
49 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide
51 <400> SEQUENCE: 3
52 Val Leu Pro Ala Leu Pro
54   1           5
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 16
58 <212> TYPE: PRT
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide

```

The type of errors shown exist throughout  
the Sequence Listing. Please check subsequent  
sequences for similar errors.



- Must explain  
genetic source -  
See error summary  
sheet item 11

## RAW SEQUENCE LISTING

DATE: 10/30/2002

PATENT APPLICATION: US/10/028,075A

TIME: 15:01:56

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

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64 <400> SEQUENCE: 4
65 Met Leu Ala Arg Arg Lys Pro Val Leu Pro Ala Leu Thr Ile Asn Pro
66   1           5           10           15
68 <210> SEQ ID NO: 5
69 <211> LENGTH: 7
70 <212> TYPE: PRT
71 <213> ORGANISM: Artificial Sequence
73 <220> FEATURE:
74 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide /
76 <400> SEQUENCE: 5
77 Met Leu Ala Arg Arg Lys Pro
78   1           5
80 <210> SEQ ID NO: 6
81 <211> LENGTH: 4
82 <212> TYPE: PRT
83 <213> ORGANISM: Artificial Sequence
85 <220> FEATURE:
86 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide /
88 <400> SEQUENCE: 6
89 Met Leu Ala Arg
90   1
92 <210> SEQ ID NO: 7
93 <211> LENGTH: 6
94 <212> TYPE: PRT
95 <213> ORGANISM: Artificial Sequence
97 <220> FEATURE:
98 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide /
100 <400> SEQUENCE: 7
101 Val Leu Pro Ala Leu Thr
102   1           5
104 <210> SEQ ID NO: 8
105 <211> LENGTH: 5
106 <212> TYPE: PRT
108 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: Description of Artificial Sequence:
113     pdb/1QMH/1QMH-A
115 <400> SEQUENCE: 8
116 Val Leu Pro Ala Leu
117   1           5
119 <210> SEQ ID NO: 9
120 <211> LENGTH: 4
121 <212> TYPE: PRT
122 <213> ORGANISM: Artificial Sequence
124 <220> FEATURE:
125 <223> OTHER INFORMATION: Description of Artificial Sequence:
126     pdb/4NOS/4NOS-A
128 <400> SEQUENCE: 9
129 Phe Pro Gly Cys

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## RAW SEQUENCE LISTING

DATE: 10/30/2002

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TIME: 15:01:56

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

```

130 1
132 <210> SEQ ID NO: 10
133 <211> LENGTH: 4
134 <212> TYPE: PRT
135 <213> ORGANISM: Artificial Sequence
137 <220> FEATURE:
138 <223> OTHER INFORMATION: Description of Artificial Sequence: Hs.297775.1
140 <400> SEQUENCE: 10
141 Pro Gly Cys Pro
142 1
144 <210> SEQ ID NO: 11
145 <211> LENGTH: 7
146 <212> TYPE: PRT
147 <213> ORGANISM: Artificial Sequence
149 <220> FEATURE:
150 <223> OTHER INFORMATION: Description of Artificial Sequence:
151     swiss/P81272/NS2B HUMAN
153 <400> SEQUENCE: 11
154 Gly Val Leu Pro Ala Val Pro
155 1 5
157 <210> SEQ ID NO: 12
158 <211> LENGTH: 6
159 <212> TYPE: PRT
160 <213> ORGANISM: Artificial Sequence
162 <220> FEATURE:
163 <223> OTHER INFORMATION: Description of Artificial Sequence:
164     swiss/P81272/NS2B HUMAN
166 <400> SEQUENCE: 12
167 Val Leu Pro Ala Val Pro
168 1 5
170 <210> SEQ ID NO: 13
171 <211> LENGTH: 4
172 <212> TYPE: PRT
173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Description of Artificial Sequence:
177     pdb/1FZV/1FZV-A
179 <400> SEQUENCE: 13
180 Pro Ala Val Pro
181 1
183 <210> SEQ ID NO: 14
184 <211> LENGTH: 9
185 <212> TYPE: PRT
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
189 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide
191 <400> SEQUENCE: 14
192 Leu Gln Gly Val Val Pro Arg Gly Val
193 1 5

```

## RAW SEQUENCE LISTING

DATE: 10/30/2002

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TIME: 15:01:56

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

195 <210> SEQ ID NO: 15  
196 <211> LENGTH: 4  
197 <212> TYPE: PRT  
198 <213> ORGANISM: Artificial Sequence  
200 <220> FEATURE:  
201 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
203 <400> SEQUENCE: 15  
204 Gly Val Val Pro  
205 1  
207 <210> SEQ ID NO: 16  
208 <211> LENGTH: 5  
209 <212> TYPE: PRT  
210 <213> ORGANISM: Artificial Sequence  
212 <220> FEATURE:  
213 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
216 <400> SEQUENCE: 16  
217 Val Pro Arg Gly Val  
218 1 5  
222 <210> SEQ ID NO: 17  
223 <211> LENGTH: 4  
224 <212> TYPE: PRT  
225 <213> ORGANISM: Artificial Sequence  
227 <220> FEATURE:  
228 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
230 <400> SEQUENCE: 17  
231 Pro Arg Gly Val  
232 1  
234 <210> SEQ ID NO: 18  
235 <211> LENGTH: 5  
236 <212> TYPE: PRT  
237 <213> ORGANISM: Artificial Sequence  
239 <220> FEATURE:  
240 <223> OTHER INFORMATION: Description of Artificial Sequence: polypeptide  
242 <400> SEQUENCE: 18  
243 Met Ala Pro Lys Lys  
244 1  
246 <210> SEQ ID NO: 19  
247 <211> LENGTH: 4  
248 <212> TYPE: PRT  
249 <213> ORGANISM: Artificial Sequence  
251 <220> FEATURE:  
252 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
254 <400> SEQUENCE: 19  
255 Leu Gln Gly Ala  
256 1  
258 <210> SEQ ID NO: 20  
259 <211> LENGTH: 10  
260 <212> TYPE: PRT  
261 <213> ORGANISM: Artificial Sequence

## RAW SEQUENCE LISTING

DATE: 10/30/2002

PATENT APPLICATION: US/10/028,075A

TIME: 15:01:56

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

263 <220> FEATURE:  
264 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
266 <400> SEQUENCE: 20  
267 Val Leu Pro Ala Leu Pro Gln Val Val Cys  
268 1 5 10  
270 <210> SEQ ID NO: 21  
271 <211> LENGTH: 6  
272 <212> TYPE: PRT  
273 <213> ORGANISM: Artificial Sequence  
276 <220> FEATURE:  
277 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
279 <400> SEQUENCE: 21  
280 Ala Leu Pro Ala Leu Pro  
281 1 5  
283 <210> SEQ ID NO: 22  
284 <211> LENGTH: 6  
285 <212> TYPE: PRT  
286 <213> ORGANISM: Artificial Sequence  
288 <220> FEATURE:  
289 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
291 <400> SEQUENCE: 22  
292 Val Ala Pro Ala Leu Pro  
293 1 5  
295 <210> SEQ ID NO: 23  
296 <211> LENGTH: 7  
297 <212> TYPE: PRT  
298 <213> ORGANISM: Artificial Sequence  
300 <220> FEATURE:  
301 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
303 <400> SEQUENCE: 23  
304 Ala Leu Pro Ala Leu Pro Gln  
305 1 5  
307 <210> SEQ ID NO: 24  
308 <211> LENGTH: 7  
309 <212> TYPE: PRT  
310 <213> ORGANISM: Artificial Sequence  
312 <220> FEATURE:  
313 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
315 <400> SEQUENCE: 24  
316 Val Leu Pro Ala Ala Pro Gln  
317 1 5  
319 <210> SEQ ID NO: 25  
320 <211> LENGTH: 7  
321 <212> TYPE: PRT  
322 <213> ORGANISM: Artificial Sequence  
324 <220> FEATURE:  
325 <223> OTHER INFORMATION: Description of Artificial Sequence: oligopeptide  
327 <400> SEQUENCE: 25  
328 Val Leu Pro Ala Leu Ala Gln

RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 10/30/2002  
PATENT APPLICATION: US/10/028,075A      TIME: 15:01:57

Input Set : A:\EP.txt  
Output Set: N:\CRF4\10302002\J028075A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:97; Xaa Pos. 2  
Seq#:98; Xaa Pos. 2

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:2; Line(s) 42  
Seq#:103; Line(s) 1376  
Seq#:118; Line(s) 1565

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/028,075A

DATE: 10/30/2002

TIME: 15:01:57

Input Set : A:\EP.txt

Output Set: N:\CRF4\10302002\J028075A.raw

L:1287 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:97  
L:1292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:97 after pos.:0  
L:1312 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:0